



OIEP

RAW SEQUENCE LISTING

DATE: 01/27/2003

PATENT APPLICATION: US/10/005,480A

TIME: 10:07:02

Input Set : D:\51158-20062.txt

Output Set: N:\CRF4\01242003\J005480A.raw

4 <110> APPLICANT: Agensys
 5 Challita-Eid, Pia M.
 6 Raitano, Arthur B.
 7 Faris, Mary
 8 Hubert, Rene S.
 9 Morrison, Karen Jane Meyrick
 10 Jakobovits, Aya
 12 <120> TITLE OF INVENTION: Nucleic Acid and Corresponding Protein
 13 Entitled 161P2F10B Useful in Treatment and Detection of
 14 Cancer
 16 <130> FILE REFERENCE: 51158-20062.00
 18 <140> CURRENT APPLICATION NUMBER: US 10/005,480A
 19 <141> CURRENT FILING DATE: 2001-11-07
 21 <160> NUMBER OF SEQ ID NOS: 765
 23 <170> SOFTWARE: FastSEQ for Windows Version 4.0

Does Not Comply
Corrected Diskette Needed

ERRORED SEQUENCES

8421 <210> SEQ ID NO: 765
 8422 <211> LENGTH: 871
 8423 <212> TYPE: PRT
 8424 <213> ORGANISM: Homo Sapien
 8426 <400> SEQUENCE: 765
 8427 Met Glu Ser Thr Leu Thr Leu Ala Thr Glu Gln Pro Val Lys Lys Asn
 8428 1 5 10 15
 8429 Thr Leu Lys Lys Tyr Lys Ile Ala Cys Ile Val Leu Leu Ala Leu Leu
 8430 20 25 30
 8431 Val Ile Met Ser Leu Gly Leu Gly Leu Gly Leu Arg Lys Leu
 8432 35 40 45
 8433 Glu Lys Gln Gly Ser Cys Arg Lys Lys Cys Phe Asp Ala Ser Phe Arg
 8434 50 55 60
 8435 Gly Leu Glu Asn Cys Arg Cys Asp Val Ala Cys Lys Asp Arg Gly Asp
 8436 65 70 75 80
 8437 Cys Cys Trp Asp Phe Glu Asp Thr Cys Val Glu Ser Thr Arg Ile Trp
 8438 85 90 95
 8439 Met Cys Asn Lys Phe Arg Cys Gly Glu Thr Arg Leu Glu Ala Ser Leu
 8440 100 105 110
 8441 Cys Ser Cys Ser Asp Asp Cys Leu Gln Lys Lys Asp Cys Ala Asp
 8442 115 120 125
 8443 Tyr Lys Ser Val Cys Gln Gly Glu Thr Ser Trp Leu Glu Glu Asn Cys
 8444 130 135 140
 8445 Asp Thr Ala Gln Gln Ser Gln Cys Pro Glu Gly Phe Asp Leu Pro Pro

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8446	145				150					155			160
8447	Val	Ile	Leu	Phe	Ser	Met	Asp	Gly	Phe	Arg	Ala	Glu	Tyr
8448					165					170			175
8449	Trp	Asp	Thr	Leu	Met	Pro	Asn	Ile	Asn	Lys	Leu	Lys	Thr
8450				180					185				190
8451	His	Ser	Lys	Tyr	Met	Arg	Ala	Met	Tyr	Pro	Thr	Lys	Thr
8452			195					200				205	
8453	His	Tyr	Thr	Ile	Val	Thr	Gly	Leu	Tyr	Pro	Glu	Ser	His
8454		210				215					220		
8455	Asn	Asn	Met	Tyr	Asp	Val	Leu	Asn	Lys	Asn	Phe	Ser	Leu
8456	225				230					235			240
8457	Glu	Gln	Asn	Pro	Ala	Trp	Trp	His	Gly	Gln	Pro	Met	Trp
8458				245					250				255
8459	Met	Tyr	Gln	Gly	Leu	Lys	Ala	Ala	Thr	Tyr	Phe	Trp	Pro
8460			260						265				270
8461	Val	Ala	Ile	Asn	Gly	Ser	Phe	Pro	Ser	Ile	Tyr	Met	Pro
8462			275					280					285
8463	Ser	Val	Pro	Phe	Glu	Glu	Arg	Ile	Ser	Thr	Leu	Leu	Lys
8464		290					295					300	
8465	Leu	Pro	Lys	Ala	Glu	Arg	Pro	Arg	Phe	Tyr	Thr	Met	Tyr
8466	305				310					315			320
8467	Pro	Asp	Ser	Ser	Gly	His	Ala	Gly	Gly	Pro	Val	Ser	Ala
8468				325					330				335
8469	Lys	Ala	Leu	Gln	Val	Val	Asp	His	Ala	Phe	Gly	Met	Leu
8470			340					345					350
8471	Leu	Lys	Gln	Arg	Asn	Ile	Leu	His	Asn	Arg	Val	Asn	Ile
8472		355					360					365	
8473	Ala	Asp	His	Gly	Met	Asp	Gln	Thr	Tyr	Cys	Asn	Lys	Met
8474		370					375					380	
8475	Thr	Asp	Tyr	Phe	Pro	Arg	Ile	Asn	Phe	Phe	Tyr	Met	Tyr
8476	385				390					395			400
8477	Ala	Pro	Arg	Ile	Arg	Ala	His	Asn	Ile	Pro	His	Asp	Phe
8478				405					410				415
8479	Asn	Ser	Glu	Glu	Ile	Val	Arg	Asn	Leu	Ser	Cys	Arg	Lys
8480			420					425					430
8481	His	Phe	Lys	Pro	Tyr	Leu	Thr	Pro	Asp	Leu	Pro	Lys	Arg
8482			435					440					445
8483	Ala	Lys	Asn	Val	Arg	Ile	Asp	Lys	Val	His	Leu	Phe	Val
8484		450					455					460	
8485	Trp	Leu	Ala	Val	Arg	Ser	Lys	Ser	Asn	Thr	Asn	Cys	Gly
8486	465				470					475			480
8487	His	Gly	Tyr	Asn	Asn	Glu	Phe	Arg	Ser	Met	Glu	Ala	Ile
8488				485					490				495
8489	His	Gly	Pro	Ser	Phe	Lys	Glu	Lys	Thr	Glu	Val	Glu	Pro
8490			500					505					510
8491	Ile	Glu	Val	Tyr	Asn	Leu	Met	Cys	Asp	Leu	Leu	Arg	Ile
8492			515					520					525
8493	Pro	Asn	Asn	Gly	Thr	His	Gly	Ser	Leu	Asn	His	Leu	Leu
8494		530					535					540	

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8495	Phe	Tyr	Glu	Pro	Ser	His	Ala	Glu	Glu	Val	Ser	Lys	Phe	Ser	Val	Cys
8496	545					550					555					560
8497	Gly	Phe	Ala	Asn	Pro	Leu	Pro	Thr	Glu	Ser	Leu	Asp	Cys	Phe	Cys	Pro
8498				565						570						575
8499	His	Leu	Gln	Asn	Ser	Thr	Gln	Leu	Glu	Gln	Val	Asn	Gln	Met	Leu	Asn
8500				580						585					590	
8501	Leu	Thr	Gln	Glu	Glu	Ile	Thr	Ala	Thr	Val	Lys	Val	Asn	Leu	Pro	Phe
8502			595					600					605			
8503	Gly	Arg	Pro	Arg	Val	Leu	Gln	Lys	Asn	Val	Asp	His	Cys	Leu	Leu	Tyr
8504		610					615					620				
8505	His	Arg	Glu	Tyr	Val	Ser	Gly	Phe	Gly	Lys	Ala	Met	Arg	Met	Pro	Met
8506	625					630					635					640
8507	Trp	Ser	Ser	Tyr	Thr	Val	Pro	Gln	Leu	Gly	Asp	Thr	Ser	Pro	Leu	Pro
8508					645					650					655	
8509	Pro	Thr	Val	Pro	Asp	Cys	Leu	Arg	Ala	Asp	Val	Arg	Val	Pro	Pro	Ser
8510				660					665					670		
8511	Glu	Ser	Gln	Lys	Cys	Ser	Phe	Tyr	Leu	Ala	Asp	Lys	Asn	Ile	Thr	His
8512			675					680					685			
8513	Gly	Phe	Leu	Tyr	Pro	Pro	Ala	Ser	Asn	Arg	Thr	Ser	Asp	Ser	Gln	Tyr
8514		690					695					700				
8515	Asp	Ala	Leu	Ile	Thr	Ser	Asn	Leu	Val	Pro	Met	Tyr	Glu	Glu	Phe	Arg
8516	705					710					715					720
8517	Lys	Met	Trp	Asp	Tyr	Phe	His	Ser	Val	Leu	Ile	Lys	His	Ala	Thr	
8518				725						730					735	
8519	Glu	Arg	Asn	Gly	Val	Asn	Val	Val	Ser	Gly	Pro	Phe	Asp	Tyr	Asn	Tyr
8520				740					745				750			
8521	Asp	Gly	His	Phe	Asp	Ala	Pro	Asp	Glu	Ile	Thr	Lys	His	Leu	Ala	Asn
8522			755					760					765			
8523	Thr	Asp	Val	Pro	Ile	Pro	Thr	His	Tyr	Phe	Val	Val	Leu	Thr	Ser	Cys
8524		770					775					780				
8525	Lys	Asn	Ser	His	Thr	Pro	Glu	Asn	Cys	Pro	Gly	Trp	Leu	Asp	Val	Leu
8526	785					790					795					800
8527	Pro	Phe	Ile	Ile	Pro	His	Arg	Pro	Thr	Asn	Val	Glu	Ser	Cys	Pro	Glu
8528					805					810					815	
8529	Gly	Lys	Pro	Glu	Ala	Leu	Trp	Val	Glu	Glu	Arg	Phe	Thr	Ala	His	Ile
8530				820						825					830	
8531	Ala	Arg	Val	Arg	Asp	Val	Glu	Leu	Leu	Thr	Gly	Leu	Asp	Phe	Tyr	Gln
8532			835					840					845			
8533	Asp	Lys	Val	Gln	Pro	Val	Ser	Glu	Ile	Leu	Gln	Leu	Lys	Thr	Tyr	Leu
8534		850					855						860			
8535	Pro	Thr	Phe	Glu	Thr	Thr	Ile									
8536	865						870									

E--> 8540 1
E--> 8543 1

VERIFICATION SUMMARY

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Input Set : D:\51158-20062.txt

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L:7450 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:742 after pos.:0
L:8410 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:763 after pos.:0
L:8540 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:765
M:332 Repeated in SeqNo=765